

0590  
0709

#10



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OIPE

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/905,558C

DATE: 07/10/2002 Pib  
TIME: 13:06:39

Input Set : A:\1016.SEQLIST.TXT

Output Set: N:\CRF3\07102002\I905558C.raw

4 <110> APPLICANT: Garnaat, Carl W.  
5 Lowe, Keith S.  
6 Roth, Bradley A.  
8 <120> TITLE OF INVENTION: ZmAxigl Polynucleotides and Methods of  
9 Use  
11 <130> FILE REFERENCE: 1016  
13 <140> CURRENT APPLICATION NUMBER: US 09/905,558C  
C--> 14 <141> CURRENT FILING DATE: 2002-06-24  
16 <150> PRIOR APPLICATION NUMBER: US 60/217,942  
17 <151> PRIOR FILING DATE: 2000-07-13  
19 <160> NUMBER OF SEQ ID NOS: 21  
21 <170> SOFTWARE: FastSEQ for Windows Version 3.0  
23 <210> SEQ ID NO: 1  
24 <211> LENGTH: 1271  
25 <212> TYPE: DNA  
26 <213> ORGANISM: Zea mays  
28 <220> FEATURE:  
29 <221> NAME/KEY: CDS  
30 <222> LOCATION: (170)...(763)  
32 <400> SEQUENCE: 1  
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34 acactagctg ctgtgggacg atcgaagtgg gtgtgtcagc tagctagctg cgccgtgacc 120  
35 acgcacatga ccgcagtgcg cgcggggctg atcaagggaa agtgatcgg atg gag ctg 178  
36 Met Glu Leu  
37 1  
39 gag ctc ggg ctc gcg ccg ccg aac ccg cat cag ccg ctg gct gcc gcc 226  
40 Glu Leu Gly Leu Ala Pro Pro Asn Pro His Gln Pro Leu Ala Ala Ala  
41 5 10 15  
43 gcc gag ttc gtc ggt ctc ctc agc agc tcg gct ggc tcg tgc ggg aac 274  
44 Ala Glu Phe Val Gly Leu Leu Ser Ser Ser Ala Gly Ser Cys Gly Asn  
45 20 25 30 35  
47 aag agg gtt ctc ggc gac gcg ttc ggg gcc gcc aag gcg gcc acg ctt 322  
48 Lys Arg Val Leu Gly Asp Ala Phe Gly Ala Ala Lys Ala Ala Thr Leu  
49 40 45 50  
51 ccg ctc ttc gtc tgc gag gat ggc gac gga ggc ggc ggc gac cgc gac 370  
52 Pro Leu Phe Val Cys Glu Asp Gly Asp Gly Gly Gly Gly Asp Arg Asp  
53 55 60 65  
55 cgc gac ggc gtc gtc gac cat gaa cag caa agc aac aat gta ccc agg 418  
56 Arg Asp Gly Val Val Asp His Glu Gln Gln Ser Asn Asn Val Pro Arg  
57 70 75 80  
59 aag aag agg ctg gtg ggg tgg ccg ccg gtg aag tgc gcg cgt agg cgt 466  
60 Lys Lys Arg Leu Val Gly Trp Pro Pro Val Lys Cys Ala Arg Arg Arg  
61 85 90 95

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63  agc tgc ggc ggc ggg tac gtg aag gtg aag ctg gaa ggg gtg ccc atc      514
64  Ser Cys Gly Gly Gly Tyr Val Lys Val Lys Leu Glu Gly Val Pro Ile
65  100                               105                               110                               115
67  ggg cgg aag gtg gac gtg tcc atc cac ggc tgc tac cag gag ctg ctc      562
68  Gly Arg Lys Val Asp Val Ser Ile His Gly Ser Tyr Gln Glu Leu Leu
69                               120                               125                               130
71  cgc acg ctc gag agc atg ttc cct tgc ggt aac caa caa gat cat gca      610
72  Arg Thr Leu Glu Ser Met Phe Pro Ser Gly Asn Gln Gln Asp His Ala
73                               135                               140                               145
75  gaa gac gag gtg gtg gtc tgc cac gag cgc cgc cgt cgc cat cct tat      658
76  Glu Asp Glu Val Val Val Ser His Glu Arg Arg Arg Arg His Pro Tyr
77                               150                               155                               160
79  gta gtc acc tac gag gac ggc gaa ggg gac tgg ttg ctc gtc gga gat      706
80  Val Val Thr Tyr Glu Asp Gly Glu Gly Asp Trp Leu Leu Val Gly Asp
81                               165                               170                               175
83  gat gtg ccg tgg gag gtc ttt gtc aag tca gtg aag cgg ctc aag ata      754
84  Asp Val Pro Trp Glu Val Phe Val Lys Ser Val Lys Arg Leu Lys Ile
85  180                               185                               190                               195
87  ctt gcg tag ccgacggctcg gcgcctcaga gacgtcgtgt ggtccgtctc      803
88  Leu Ala *
91  accaggatcg gacgagtgtg gtactcctgg gcgtcatctg cgtaataacg ttgtttctgt      863
92  cctgtgtgcc cgtagcagta cgtactgtcc tataagtaagc tagctttatg ggggtgcttca      923
93  gctttcagag catgacgaaa gcaactgatta gctgctgtca tcacatttgg ttcgtctttg      983
94  tgctgtacgg tatcgctggc gtcagtgtcg cggcagccta ggtgatctaa gcatacttac      1043
95  tatctcaagt tacttttggg ttcctgagct tgcattgtaa ttcataatacc gtatacgtgt      1103
96  gtgactcagg ggcgaagctg ccttaaggca caggggtcac cggaccgat ggaatttatc      1163
97  aaatccagtg taaaatacta ttttaactg ttcataata tatttgattt caataaaaaa      1223
98  aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa      1271
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101 <211> LENGTH: 197
102 <212> TYPE: PRT
103 <213> ORGANISM: Zea mays
105 <400> SEQUENCE: 2
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108 Ala Ala Ala Ala Glu Phe Val Gly Leu Leu Ser Ser Ser Ala Gly Ser
109 20 25 30
110 Cys Gly Asn Lys Arg Val Leu Gly Asp Ala Phe Gly Ala Ala Lys Ala
111 35 40 45
112 Ala Thr Leu Pro Leu Phe Val Cys Glu Asp Gly Asp Gly Gly Gly Gly
113 50 55 60
114 Asp Arg Asp Arg Asp Gly Val Val Asp His Glu Gln Gln Ser Asn Asn
115 65 70 75 80
116 Val Pro Arg Lys Lys Arg Leu Val Gly Trp Pro Pro Val Lys Cys Ala
117 85 90 95
118 Arg Arg Arg Ser Cys Gly Gly Gly Tyr Val Lys Val Lys Leu Glu Gly
119 100 105 110
120 Val Pro Ile Gly Arg Lys Val Asp Val Ser Ile His Gly Ser Tyr Gln
121 115 120 125

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122  Glu Leu Leu Arg Thr Leu Glu Ser Met Phe Pro Ser Gly Asn Gln Gln
123      130                      135                      140
124  Asp His Ala Glu Asp Glu Val Val Val Ser His Glu Arg Arg Arg Arg
125      145                      150                      155                      160
126  His Pro Tyr Val Val Thr Tyr Glu Asp Gly Glu Gly Asp Trp Leu Leu
127      165                      170                      175
128  Val Gly Asp Asp Val Pro Trp Glu Val Phe Val Lys Ser Val Lys Arg
129      180                      185                      190
130  Leu Lys Ile Leu Ala
131      195
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134 <211> LENGTH: 1310
135 <212> TYPE: DNA
136 <213> ORGANISM: Zea mays
138 <400> SEQUENCE: 3
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140  tgttcttatt cagactaccg ttcgagtgac tgcattggcg acatctttct gcatcgactt      120
141  tgtacggcta catcgaacat atacacgaga tgtctcgtgt gaatagagtc actaatgcct      180
142  taagcatcgg ttactccgta gggtagattc tgtttcttct atttgtgcat atttttattg      240
143  ttgtttactg attatacgag tagttataca tacatgcaca tacatatcat cacatatatc      300
144  acaatatatt tctaaattaa attaaaacta aaaatgacta aattttctaac accaacgaca      360
145  ttgtaatggt ttctccaaca actttaccta ttctacattg ttctatttcg aattttcactc      420
146  tataaacaac atagtctaca atggaaaaca gtgctttgta cgactatata cgcgatgtgt      480
147  ggctacaaca taagacaata tagtcgtttg aagattgaac ctatatatcg gtacgggtta      540
148  tccgtctatg tacgtgggca tgacgaacac ccgtgataac gaaggattaa cgtgcacaat      600
149  cataaatcca aagtaggagc ggtgcatgat gagaatcgct ctcagtactc gacataatga      660
150  accttacgag gtacaacagg caggcaggca gggaccagg gccgccttta ttccaggctc      720
151  gctggcccca cgggcgtgct gcgtgcacga agggcactac cccaacctct caccgaaaaa      780
152  ccgcgctgga tcggcaaata aaacgaggtg gtgccccgtg cccactctcc acgtccacgg      840
153  caccatccct ctgcagccgc tcaccagcca tgccgtgtcg cggaaacggc caaccacccc      900
154  caaccactc acgaaacccc gtcccggcgc tgcccgtgtc ggtccgcgct cggcaacgag      960
155  gcggcccgcg ctgctgagtc ccctggacac ccgacaccct gtcggccctt tgtttattca      1020
156  tcccgaaatc tcatctgccc ccacggccga ctgctgctgc ccgcccggat atatataccc      1080
157  atcgttatcg atcgatcgat cgcgtcactc acgggtagct catggtcgag cgtagcatgc      1140
158  aggaacttat ttgccgtgcg ctcccaggtc tccgctcgcg tgccttccag tctgtctcac      1200
159  actagctgct gtgggacgat cgaagtgggt gtgtcagcta gctagctgcg ccgtgaccac      1260
160  gcacatgacc gcagtgcgcg cggggtgat caagggaag tgatcggatg      1310
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163 <211> LENGTH: 1310
164 <212> TYPE: DNA
165 <213> ORGANISM: Zea mays
167 <400> SEQUENCE: 4
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169  tgttcttatt cagactaccg ttcgagtgac tgcattggcg acatctttct gcatcgactt      120
170  tgtacggcta catcgaacat atacacgaga tgtctcgtgt gaatagagtc actaatgcct      180
171  taagcatcgg ttactccgta gggtagattc tgtttcttct atttgtgcat atttttattg      240
172  ttgtttactg attatacgag tagttataca tacatgcaca tacatatcat cacatatatc      300
173  acaatatatt tctaaattaa attaaaacta aaaatgacta aattttctaac accaacgaca      360
174  ttgtaatggt ttctccaaca actttaccta ttctacattg ttctatttcg aattttcactc      420

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175 tataaacaac atagtctaca atggaaaaca gtgctttgta cgactatata cgcgatgtgt      480
176 ggctacaaca taagacaata tagtcgtttg aagattgaac ctatatatcg gtacggttaa      540
177 tccgtctatg tacgtgggca tgacgaacac ccgatgataac gaaggattaa cgtgcacaat      600
178 cataaatcca aagtaggagc ggtgcatgat gagaatcgct ctcagtactc gacataatga      660
179 accttacgag gtacaacagg caggcaggca gggaccaggg gccgccttta tttcaggctc      720
180 gctggcccca cgggcgtgct gcgtgcacga agggcactac cccaacctct caccgaaaaa      780
181 ccgcgctgga tcggcaaata aaacgagggt gtgccccgtg cccactctcc acgtccacgg      840
182 caccatccct ctgcagccgc tcaccagcca tgcccggtgc cggaacggca caaccacccc      900
183 caaccactc acgaaacccc gtcccggccg tgcccgtgtc ggtccgcgct cggcaacgag      960
184 gcggcccgcg ctgctgagtc ccctggacac ccgacaccct gtcggccctt tgtttattca    1020
185 tcccgaatac tcatctgccc ccacggccga ctgcgctgcg ccgcccgat atatatacc      1080
186 atcgttatcg atcgatcgat cgcgtcactc acgggtagct catggtcgag cgtagcatgc      1140
187 aggaacttat ttgcgctgcg ctcccaggtc tccgctcgcg tgccctccag tctgtctcac      1200
188 actagctgct gtgggacgat cgaagtgggt gtgtcagcta gctagctgcg ccgtgaccac      1260
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192 <211> LENGTH: 3123
193 <212> TYPE: DNA
194 <213> ORGANISM: Zea mays
196 <400> SEQUENCE: 5
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199 tgtacggcta catogaacat atacacgaga tgtctcgtgt gaatagagtc actaatgcct      180
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201 ttgtttactg attatacgag tagttataca tacatgcaca tacatatcat cacataatc      300
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204 tataaacaac atagtctaca atggaaaaca gtgctttgta cgactatata cgcgatgtgt      480
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206 tccgtctatg tacgtgggca tgacgaacac ccgatgataac gaaggattaa cgtgcacaat      600
207 cataaatcca aagtaggagc ggtgcatgat gagaatcgct ctcagtactc gacataatga      660
208 accttacgag gtacaacagg caggcaggca gggaccaggg gccgccttta tttcaggctc      720
209 gctggcccca cgggcgtgct gcgtgcacga agggcactac cccaacctct caccgaaaaa      780
210 ccgcgctgga tcggcaaata aaacgagggt gtgccccgtg cccactctcc acgtccacgg      840
211 caccatccct ctgcagccgc tcaccagcca tgcccggtgc cggaacggca caaccacccc      900
212 caaccactc acgaaacccc gtcccggccg tgcccgtgtc ggtccgcgct cggcaacgag      960
213 gcggcccgcg ctgctgagtc ccctggacac ccgacaccct gtcggccctt tgtttattca    1020
214 tcccgaatac tcatctgccc ccacggccga ctgcgctgcg ccgcccgat atatatacc      1080
215 atcgttatcg atcgatcgat cgcgtcactc acgggtagct catggtcgag cgtagcatgc      1140
216 aggaacttat ttgcgctgcg ctcccaggtc tccgctcgcg tgccctccag tctgtctcac      1200
217 actagctgct gtgggacgat cgaagtgggt gtgtcagcta gctagctgcg ccgtgaccac      1260
218 gcacatgacc gcagtgcgcg cggggctgat caagggaaag tgatcggatg gagctggagc      1320
219 tcgggctcgc gccgcgaac ccgcatcagc cgctggctgc gccgcgcgag ttctgctggc      1380
220 tcctcagcag ctgcgctggc tcgtgcggga acaagagggt tctcggcgac gcgttcgggg      1440
221 ccgccaaggc ggccaacgct ccgctcttcg tctgcgagga tggcgacgga ggcggcgcg      1500
222 accgcgaccg cgacggcgct gtcgaccatg aacagcaaag caacaagtga gttgtggtta      1560
223 aaaataccga ccacgtgcgt acagggaggg tcttattata cccaaatccg atccgtgggt      1620
224 tgtgtagtgt acccaggaag aagaggctgg tggggtgccc gccggtgaag tgcgcgcgta      1680
225 ggcgtagctg cggcggcggg tacgtgaagg tgaagctgga aggggtgccc atcgggcgga      1740

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227 tcccttcggg taaccaacaa ggtgcgtacg ttcccggggc gcggcgagcc ggccggcgac 1860
228 cggcggtgct gcggacgatg cctttctttc actgataatc atctgccgcc atcgttcttg 1920
229 tcccgcacag tgcccttgct tcccgttctg ctcccggcac ttaacttggt cgcataact 1980
230 attcctgtaa cctctggcag atcatgcaga agacgaggtg gtggtctcgc acgagcgccg 2040
231 ccgtcgccat ccttatgtag tcacctacga ggacggcgaa ggggactggt tgctcgtcgg 2100
232 agatgatgtg ccgtgggagt acgtatcagt cactactact gtcgtctgta tgactgtatc 2160
233 gatggtgacg gcaacaatat aatccaatta attattcagc gaacttaaaa acgacgttga 2220
234 tttccttgca gggctcttgt caagtcagtg aagcggtcga agatacttgc gtagccgacg 2280
235 gtcggcgccct cagagacgct gtgtgggtccg tctcaccagg atcggagcag tgtagtactc 2340
236 ctgggctgca tctgcgtaat aacgttggtt ctgtcctgtg tgcccgtagc agtacgtact 2400
237 gtcctatagt aagctagctt tatggggtgc ttacgctttc agagcatgac gaaagcactg 2460
238 attagctgct gtcacacat ttggttcgtc tttgtgtcgt acggtatcgc tggcgtcagt 2520
239 gtcgcggcag cctaggtgat ctaagcatac ttactatctc aagttacttt tggtttcctg 2580
240 agcttgcatg gtaattcata taccgtatac gtgtgtgact caggggcgaa gctgccttaa 2640
241 ggcacagggg tcaccggacc cgatggaatt tatcaaatcc agtgtaaaat actatttaac 2700
242 actgttcata aatataattt atttcaataa ttcattggagc tgaccttggt gatccatttt 2760
243 ctgtcttcgc ctctggtgtg actagtattt tggtttgact tttcactctg tataagatat 2820
244 atattatacc agcgagttaa tcgactgcta gttttacaag aggcttaact ctttcaattg 2880
245 cttattttta ttgcaacaac acactcctcc gttgttggtg tattagatgt ggttctgaat 2940
246 gtaaatgtca ttataggata taaatgtagt gtttcctagt tttaccctag ctttcgcatg 3000
247 catagtggga aagtgtacta actctcctca tgcagaaaga ggtgtggtat acctaacaaa 3060
248 atcatacatc actactaatc tacggataat atatataaac cgtagcgaca cacgagtgt 3120
249 tag 3123
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252 <211> LENGTH: 28
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254 <213> ORGANISM: Zea mays
256 <400> SEQUENCE: 6
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260 <211> LENGTH: 28
261 <212> TYPE: DNA
262 <213> ORGANISM: Artificial Sequence
264 <220> FEATURE:
265 <223> OTHER INFORMATION: Zea mays
267 <400> SEQUENCE: 7
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270 <210> SEQ ID NO: 8
271 <211> LENGTH: 29
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273 <213> ORGANISM: Zea mays
275 <400> SEQUENCE: 8
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278 <210> SEQ ID NO: 9
279 <211> LENGTH: 29
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281 <213> ORGANISM: Zea mays
283 <400> SEQUENCE: 9

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RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 07/10/2002  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:20; N Pos. 236,242,257,276,420,457,463,470,486,508,540,554,580,586,596  
Seq#:21; Xaa Pos. 5,6,18,19,22,33,45,47,51,54,55,56,57,61,62,63,65

## VERIFICATION SUMMARY

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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:461 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:180  
L:462 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:240  
L:464 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:360  
L:465 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:420  
L:466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:480  
L:467 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:540  
L:481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0  
L:483 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:16  
L:485 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:32  
L:487 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:48  
L:489 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:64